GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:06:08; Search time 2533.16

Seconds

(without alignments)
5092.486 Million cell

updates/sec

Title: US-09-300-482-225

Perfect score: 316 Sequence: 1

gataaggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 45554873 seqs, 20411521753 residues

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Pending_Patents_NA_Main:*

1: /cgn2_6/ptodata/1/pna/PCTUS1_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/PCTUS2_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US06_COMB.seq:*

5: /cqn2 6/ptodata/1/pna/US07 COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US080_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US081_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/US082_COMB.seq:*

9: /cgn2_6/ptodata/1/pna/US083_COMB.seq:*

10: /cgn2_6/ptodata/1/pna/US084_COMB.seq:*

11: /cgn2_6/ptodata/1/pna/US085_COMB.seq:*

12: /cgn2_6/ptodata/1/pna/US086_COMB.seq:*

13: /cgn2_6/ptodata/1/pna/US087_COMB.seq:*